Appl. No.

10/786,518

Filed

February 4, 2004

## AMENDMENTS TO THE SPECIFICATION

Please replace Tables 2 through 10, which are located between paragraphs [0061] and [0062] of the specification, with the following paragraphs:

Table 2: CDH23 primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
CDH23-1F	CACTGTGCTATACCCAGGATAGGACAATGTTA	1	32	6886 bp	1F/1R	2 to 3
CDH23-1R	TCAGGTGGAAGATGACCTCAACCTGTAAGATC	2	32	·		
CDH23-2F	GATACCATCATGACACTGTGACAAGT	3	28	1426 bp	2F/2R	4 to 6
CDH23-2R	GACTCTTCACCTACACCATGGTGGTCTG	4	28			
			<b></b>			
CDH23-3F	TATGTATTCTTCACACTAACCCTGTGAGATATG	5	33	12362 bp	3F/3R	7 to 9
CDH23-3R	TAGCCCTCAGAGCCTGAGATGCCTACTGGCTC	6	32			
CDH23-4F	TGAGTCTTTAATGCCCAGAGAGGAG	Z	25	2604 bp	4F/4R	10 to 11
CDH23-4R	TGAGATGGAGTCTTACTCTTGTTGC	8	25	2004 bp	41741	10 10 11
CDI 123-41	TONONTOGRATETTACTETTACTO	<u> </u>	-23-			
CDH23-5F	CCAGAAGCTATGGCCCATCAGAGG	9	24	3425 bp	5F/5R	12 to 13
CDH23-5R	GCAACCAAGAGTACTGACAGATACA	10	25			
·						
CDH23-6F1	TGTAGGTAGAAGGCGTGCAGGAGCCAGCAGTCGC	11	34	6878bp	6F/6R	14 to 16
CDH23-6R1	GGTTCGAGTGTTTGCTGCTCAGCCTTCCGAGTAT	12	34			
			<u></u>			ļ
CDH23-6F1b	CCAAAGGAGACGTGCGAGAGGAACAT	<u>13</u>	26	4601 bp	6F1b/6R1b	14 to 16
CDH23-6R1b	TTCCTGAGTAGCCCAGAGTGTCAGG	14	25			
CDH23-7F	ACCTCAGTCGAGATGTTGAGGCTCCAGGTGTTC	<u>15</u>	33	13282 bp	7F/7R	17 to 21
CDH23-7R	CTATTGCAAGAGCCAGCTCAGAGGGACACAGA	16	32	13202 50	717713	17 (0 21
051120 111		<del>                                     </del>				
CDH23-8F	GAGGGTTTGATGAGGAGGAACCCAGTCTCCAA	17	32	12314 bp	8F/8R	22 to 27
CDH23-8R	ATTAACTCGCTGGCTCTAGGATTTCAGTAAGAG	<u>18</u>	33			
CDH23-9F	GTAGGATGCGTGAAGGGAAGGAACT	<u>19</u>	30	8499 bp	9F/9R	28 to 31
CDH23-9R	GTGCACACAGAAGGAGCTCAACCAATGTTGG	20	31			
001100 405	0774700000404040404040404040	24		7000 h -	405/405	20.4- 20
CDH23-10F CDH23-10R	GTTATGCCGGACAGAGGAAGTGACATGGAGGT CAAGGATTCGCCTGCTGTGTGGAATTCCATTC	<u>21</u> <u>22</u>	32 32	7903 bp	10F/10R	32 to 36
CDH23-TUR	CAAGGATTCGCCTGCTGTGTGGAATTCCATTC		32			
CDH23-11F	GAGTCACATGGAGTGAGTTCAGCCCAGGAGAA	23	32	11691 bp	11F/11R	37 to 43
CDH23-11R	ACAATGACCACGACTGTCTCTTCCAACCAGAC	24	32	11001 55	71177711	01 10 10
CDH23-12F3a	TTATGACTTGCTTCTGATCTTCCTTTCTGATG	<u>25</u>	32	7912bp	12F3a/12R3a	44 to 46
CDH23-12R3a	TTTGTAAAACTAGATAATTACACTACCGACTG	<u>26</u>	32			
						-
CDH23-12F4	ACACAGAGGTGCAGAGAGGTGACATAACTTCC	27	32	6815bp	12F4/12R6	47 to 53
CDH23-12R6	TAGCACAGCCCATATAGTAACCACTGTTCAATAC	<u>28</u>	34	<del></del>		
CDH23-13F	CTTCCACACCCATCATCTCTTCCCCCCCTCCCA	20	32	12462 ha	13F/13R	53 to 60
CDH23-13F CDH23-13R	CTTGGACACCCATGATGTCTTGGGGGGTGGGA GTGACCCTCCTTACCTTGTCCTTAGATGCTTAACATT	<u>29</u> 30	37	12462 bp	135/135	53 to 68
ODDZ3-13K	GIGACCCICCITACCITGICCITAGAIGCITAACATI	<u> </u>	<u> </u>		l	

Table 3: GJB2 primers

primer name	primer sequence	SEQ ID bases proc	duct size primer pair exons

		NO:				
GJB2-1F	AACCTTAGTCCTTGGCACATTGTTGAA	31	27	6478 bp	1F/1R2	1 to 2
GJB2-1R2	AACACCACATTGTCCATAGACTGATATG	<u>32</u>	28			
GJB2-1F2	AGTCAATGCTAATAATGGTGGCAATCACG	33	29	7156 bp	1F2/1R2	2
GJB2-1R2	AACACCACATTGTCCATAGACTGATATG	34	28			

Table 4: GJB6 primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
GJB6-1F	TATGAGAAGGCTGGATCACCCAGAAAGACTG	35	31	11,112 bp	1F/1R	ali 4 exons
GJB6-1R	TGAGGACATCATCCTAGTGTCGTACAAGTGG	<u>36</u>	31			
GJB6-2F-1	TGTGTTCCTGGATTAATGCAAACAGC	37	26	2361 bp	2F-1/2R-2	all 4
GJB6-2R-2	GGACATCATCCTAGTGTCGTACAAGT	38	26			
GJB6-2F-2	AGCCAATCTGGTGTAATGGATCAGAC	39	26	2383 bp	2F-2/2R-1	all 4 exons
GJB6-2R-1	AGTGCTCTGTAGGCTGCTAAACTTAG	<u>40</u>	26			

Table 5: KCNE1 primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
KCNE-1F	GAAAGAGGCATGGAGAGTGAT	41	21	1719 bp	1F/1R1	1 to 2
KCNE-1R1	CTGAAGCTCACTGACGTCTGT	42	20			
KCNE-1F1	CATGGATACCAAGAGACAACT	43	21	1724 bp	1F1/1R	
KCNE-1R	AGGATCACCTTCCTTGATTC	44	20			
KCNE-2F	TCCATTAAGGAAGGACCTTG	45	20	437bp	2F/2R	3
KCNE-2R	TAAACATTCAGCGAATGCAG	<u>46</u>	20			
KCNE-3F1	AACCAGTCTGACTAGTCTTGCATAAGCT	47	28	4893 bp	3F1/3R2	4
KCNE-3R2	GAGTCTGTTATGCTTCTGTCAGGTGT	48	28			

Table 6: KCNQ1 primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
KNQ1-1F1	GGTAAATGCACACTGGAACG	49	20	1168bp	1F1/1R1	1
KNQ1-1R1	AGGATTCACACCTGGACTAC	50	20			
KNQ1-2F	ATCCACGTGGCAGCATGTGTTG	<u>51</u>	22	564bp	2F/2R	2
KNQ1-2R	CTTTCAGACCACCAGCTCCAGGTT	<u>52</u>	24			
KNQ1-3F	ATGAGCTGAAGCTGCTCAGCCTTC	<u>53</u>	24	2709bp	3F/3R	3 to 6
KNQ1-3R	TCCAAGCACAGGTTTGTGGACAG	<u>54</u>	23			
KNQ1-4F	GCTCTGTTCCTGGTGCTTTCGCCGAGT	<u>55</u>	27	5779bp	4F/4R1	7 to 10
KNQ1-4R1	GACAGGTCTGCCATCCAATCGTCAGGT	<u>56</u>	27	6183 bp		
KNQ1-5F1	GACACTGAGGTGTCAGGCACTT	<u>57</u>	22	532bp	5F1/5R1	11
KNQ1-5R1	AGGATCATGTTCCCAGGCTCA	<u>58</u>	21			

KNQ1-6F	TTGCTATGGCTGCCATGTGTCAGCAGCATAG	<u>59</u>	31	9883bp	6F/6R	12 to 15
KNQ1-6R	TCTGCCACCCCCACTCAGGACACAGCCAG	<u>60</u>	30			
KNQ1-7F	TTGCAGACATAGGGTGCACACGTGC	<u>61</u>	25	1589BP	7F/7R	16
KNQ1-7R	AACAGGAGCGACGTCGCTAAGCTAG	<u>62</u>	25			

Table 7: MYO7A primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
MYO7A-1F	AGCACATCAGTGATTAAGTCAGG	<u>63</u>	23	822 bp	1F/1R	1
MY07A-1R	GATTCGATGGACAACATGCTCCT	<u>64</u>	23			
MYO7A-2F	TECCOATCTCTCAATCACACTC	<u>65</u>	22	424 ba	25/25	
	TTGGGAATCTCTGAATGACAGTG	66	23	434 bp	2F/2R	2
MYO7A-2R	GGTTTGGAAGCCTAGGCAGGAA	00	22			
MYO7A-3F	GAGAGGCCTTGGCTCTCTGA	<u>67</u>	22	628 bp	3F/3R	3
MYO7A-3R	TCTCTAACACCATGCAGAGTGG	<u>68</u>	22			
MYO7A-4F8	CTCATCTCACATTCCTCCTACT	69	23	206255	4E9/4D9	
	CTGATGTCCAGATTCCTGCTAGT			2863bp	4F8/4R8	4
MYO7A-4R8	ACCTCCAGCATTTATTCATGCCATG	<u>70</u>	25			
MYO7A-5F	AGAAGGAAATCTAGGCTTAGAGACTCCACCTCCC	<u>71</u>	34	7707 bp	5F/5R	5 to 14
MYO7A-5R	GCATATGATTCCACTTATATGAGGTACCTAGAAT	<u>72</u>	34			
MYO7A-6F	TGGATGTGGTGGAACTAGGTGG	<u>73</u>	22	400 bp	6F/6R	15
MYO7A-6R	AACCGATCCCTGACCGGTTCTG	74	22	488 bp	OF/OR	15
WITO/A-OIX	AACCGATCCCTGACCGGTTCTG	1-3	22			
MYO7A-7F1a	AGAGGTGGTAACTTTGGAAGTCCTGG	<u>75</u>	26	7573bp	7F1a/7R1a	16 to 21
MYO7A-7R1a	GGTATGTGCACTCCTCAGAGCAGGCATA	<u>76</u>	28			
MYO7A-7F1d	TGGTCAGATGGATAGATGGCATCACCTC	77	20	4402 5-	7544/704-0	16 to 18
		78	28	4102 bp	7F1d/7R1a2	10 10 10
MYO7A-7R1a2	ATCACATCTTGCTGATGAGGAAATGCAGG	<u>/0</u>	29			
MYO7A-7F1e	TCACAGTCTGGTGGCATAGTACCTAAATTG	<u>79</u>	30	4128 bp	7F1e/7R1a1	16 to 18
MYO7A-7R1a1	CTCCCAGGTTGTAGATGATCTCAAACAC	<u>80</u>	28	·	-	
MAYO7A 7521-	TOCACOTOCTOCTOCOC	01	20	504 h	704 - 7004 -	24
MYO7A-7F21a	TGCAGCTCCTGATCTAGGAT	<u>81</u>	20	591 bp	7F21a/7R21a	21
MYO7A-7R21a	AGAGCAGGCATAACTGCAG	<u>82</u>	21			
MYO7A-7F21b	ATTAGAGATCTCAGACAGGGTG	<u>83</u>	22	898bp	7F21b/7R21b	21
MYO7A-7R21b	AACTGGGCATGACTTTGATAGG	<u>84</u>	22			
10/074 750	1007010701070700011707070	05		20041		
MYO7A-7F2a	ACCTCAGTCACTCTTGGGAATCTCTG	<u>85</u>	26	3361bp	7F2a/7R2a	22 to 26
MYO7A-7R2a	TAGAAGTGTATTCCCTCTCAGCTGTG	<u>86</u>	26			
MYO7A-8F	TGCAGGGTATCGAGGAGGTGGC	<u>87</u>	22	620 bp	8F/8R	27
MYO7A-8R	TGCAATATCTCCAAGGGATGCC	<u>88</u>	22			
10/074 054	00000077440747701010177	00		44 ===:	0.51/0.55	20 1 2=
MYO7A-9F1	GGCCCCTTAAGTATTCACACATTACAGAAATA	89	32	11,772bp	9F1/9R3	28 to 35
MYO7A-9R3	GTTGAAACTTGATCTCCCAGTGTTGGCAGTGG	90	32			
MYO7A-10F	CGAGGTGGAAGGAGTCTGGGAGGCCCGCTCACAA	91	34	8018 bp	10F/10R	36 to 44
			<del>- '</del>	77.70		33 10 44

MYO7A-10R	AGACACATAATAGAGGCTCAACATGCAAGCTTCC	<u>92</u>	34			
MYO7A-11F	GGCCATGCACTCCAACTGCCAACTGCTGAGTCT	93	33	4555 bp	11F/11R	45 to 49
MYO7A-11R	TCACCTCCCAGCCTGATGTCCAGCACTTCCTCC	<u>94</u>	33			

Table 8: OTOF primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
OTOF-1F	TGGTAGCACATAAGCCTCTG	95	20	1001	1F/1R	1
OTOF-1R	ATCACAATGGCCAGTCAGTC	<u>96</u>	20			
OTOF-2F	TCCTAACATGGAACTCATGG	97	20	451	2F/2R	2
OTOF-2R	TTACCACCTCCTTCAGGAAG	98	20			
OTOF 25	COAACATOTOTCACCACCAT	99		700	25/20	
OTOF-3F OTOF-3R	CCAACATCTCTGAGCACCAT TGAGTGTCTGAGATCAGGC	100	20 19	786	3F/3R	3
OTOF-4F	ACAAACAACCATCCACAGTGGG	101	22	3197	4F/4R	4 to 5
OTOF-4R	TCTGAGAAAGGCAGGAGATCTAG	<u>102</u>	23			
OTOF-5F	AAAGACAAGTCAGGCTTTGAGCAC	103	24	2937	5F/5R	6 to 8
OTOF-5R	TATGAAGTCCAATACTGAACATG	<u>104</u>	23			
OTOF-6F	TGTGGTAGTGCATGCCTGTAATCC	105	24	6513	6F/6R	9 to 11
OTOF-6R	ATGGCTGTGTGTACTAACAGTCGC	106	24			
0705.754		407				
OTOF-7F1a OTOF-7R1a	AGCTCCAGAGGACCTCAGACTCTATC TGAGGTATGACTCCTCAGGTAGACAG	107 108	26 26	4152	7F1a/7R1a	12 to 25
0101-71(14	TORGOTATOROTOROTAGOTAGOTA	122	20			
OTOF-7F2a	CCTGCTTCCATGGATATCCAGGCT	<u>109</u>	24	5373	7F2a/7R2a	16 to 25
OTOF-7R2a	CTCAGTCTGTAGGAGACAGGAGGTGA	<u>110</u>	26			
OTOF-7F2e	CTGTGGAGATCGTAGACACCTCCAA	<u>111</u>	25	1791	7F2e/7R2e	16 to 18
OTOF-7R2e	ACTAGAGGTGGCTCCTGTCCTTGTC	<u>112</u>	25			
OTOF-7F2f	TAACTACACGCTGCTGGATGAGCATC	<u>113</u>	26	1784	7F2f/7R2d	16 to 18
OTOF-7R2d	AGACCAGCTTTGTGTTTCCAGGGAAG	114	27	1704	/ FZI/ RZU	10 10 10
			-			
OTOF-7F2e OTOF-7R2i	CTGTGGAGATCGTAGACACCTCCAA	115	25	3315	7F2e/7R2i	16 to 20
OTOF-7R21	CTCTGTAGATTCTTCCTCATCTGCCC TAACTACACGCTGCTGGATGAGCATC	116 117	26 26	3404	7F2f/7R2i	16 to 20
				0,01	71 2071 (21	70 10 20
OTOF-7F2m	TGATCAA'CAGGGAGGAGGCATTT	<u>118</u>	23	955	7F2m/7R2m	19 to 20
OTOF-7R2m	CTGCCCCTCCAGCACCTTA	<u>119</u>	20			
OTOF-7F2n	CCTAGCGAGAGCTCCCAG	<u>120</u>	18	542	7F2n/7R2n	19 to 20
OTOF-7R2n	GACAGCTCGGGCCATGAC	<u>121</u>	18			
OTOF-7F3f1	TGGGCAGATGAGGAAGAATCTACAGAGC	122	28	2838	7F3f1/7R3a1	21 to 25
OTOF-7R3a1	TTACCACAGCGCCATGAGTTGTTGTAAG	123	28	2000	, 1 3/1// (Jal	21 10 23
OTOF-7R3b1	ACATGAGGTCCTCCTACCTCTAGTCCAG	<u>124</u>	28	2697	7F3f1/7R3b1	21 to 25
OTOF-7F-A	CTGTGGAGATCGTAGACACCTCCAACCCTGAGCT	<u>125</u>	34	16 256	75 A/7D A	16 to 20
OTOF-/F-A	O TO TO GONGATO GIAGA TAGA TAGA TAGA TAGA TAGA TAGA TAG	152	<b>J4</b>	16,256	7F-A/7R-A	16 to 39

OTOF-7R-A	CAGATAGCCTCTCTACCTCACTGGGATTTGGACA	<u>126</u>	34			
OTOF-8F5	TAAGGACCAAACGAGATCACAGGTGTGGA	<u>127</u>	29	10127	8F5/8R6	26 to 39
OTOF-8R6	AGCCTCTCTACCTCACTGGGATTTGGACA	<u>128</u>	29			
OTOF-8R7	CGAGTCACTAGAAGTAGGATCTTGGTTTGT	<u>129</u>	30	10181	8F5/8R7	26 to 39
OTOF-8R4	GGTTTGTTCTACCTCACTGGGATTTGGACA	<u>130</u>	30	10128	8F5/8R4	26 to 39
OTOF-9F1	GTAGACAGGTGATGGCATAGAGGCTTCT	<u>131</u>	28	7106	9F1/9R1	40 to 47
OTOF-9R1	TGGTACTGAATCTGCCAGCCTAGAGAAC	<u>132</u>	28			
OTOF-9F2	AGGCACTTCCCAGAGAAGCAGAGAATTG	<u>133</u>	28	7759	9F2/9R9	40 to 47
OTOF-9R9	TGTGGCTGAATCTCTTTAAAGAGGTCAGG	<u>134</u>	29			

Table 9: SLC26A4 sequences

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
SLC-1F	* TCAGAGAATTTGCATCAGGGTTCTC	135	25	3665	1F/1R	1 to 3
SLC-1R	TAAGCAACCATCTGTCACAGACC	136	23			
SLC-2F2	TGGAACCATTGTAAGTTGAGGACTT	137	25	3225	2F2/2R4	4 to 6
SLC-2R4	GAGATGAGGTCTCACGTCTCAAACT	<u>138</u>	25			
SLC-3F	ATCAACTGGGAGTTTCAGGTTTATCAGCC	139	29	7618	3F/3R	7 to 10
SLC-3R	AAGGCAAATTGTCCTGCTAAGCTCGGTG	140	28			
SLC-4F	AATGAGACCATGTGCTACAAGTACGAAGTG	141	30	11306	4F/4R	11 to 18
SLC-4R	TTTGTTCACTCTTACCTAGGTGAGAGCCTG	142	30		-	
SLC-5F4	GATCGTCCACAAGGTTGACTACGACCAGT	143	28	9069	5F4/5R6	19 to 21
SLC-5R6	TCATTGATTCTCACCTCACAGATCTAAGC	144	29			

Table 10: USH2A sequences

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
USH2A-1F	TAGGATAAGGTGTACTGCTACTT	145	23	5085	1F/1R	1 to 3
USH2A-1R	GAAGACAAATCCTTGTGTTTAACCA	<u>146</u>	25			
USH2A-2F	AACACATGGAGATATCACTGAGC	147	23	699	2F/2R	4
USH2A-2R	CCTAAATCCAATGACAAGTGTCCTT	148	25			
USH2A-3F1	CTTAAGTCCTACAGTGTCCATGGAGATA	149	28	7298	3F1/3R1	5 to 9
USH2A-3R1	CATCAGTGATGTGTTAAAGGTTATATTC	<u>150</u>	28			
USH2A-4F	TCACTGATATGTGCTTTACTTCTGG	<u>151</u>	25	3302	4F/4R	10 to 11
USH2A-4R	AGGATTTCCTGGCAAATGCAGTCTTC	<u>152</u>	26			
USH2A-5F	GTCTTGTACCTAATGAGCAAATTATCT	153	27	4954	5F/5R	12 to 13
USH2A-5R	GCATTGTATGGATATTCAACTCAAATT	154	27			
USH2A-6F1	GAATTAGTGCCTTGGTAGA	<u>155</u>	19	378	6F1/6R	14
USH2A-6F2	GTATTGGGAATTAGTGCCTT	<u>156</u>	20	386	6F2/6R	14
USH2A-6R	CAGAAGTTATTGCTTTGCAACT	157	22			
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USH2A-7F	CTCTACAATGCTATTGGTAGGTGTAACTTA	<u>158</u>	30	10458	7F/7R	15 to 16
USH2A-7R	CACAACAGCATTTATCCTCAATGTCAAAGA	<u>159</u>	30			
USH2A-8F	AGCAGTTAGCAATGATTCTTCACCAACTTGTG	<u>160</u>	32	10312	8F/8R	17 to 20
USH2A-8R	CCTGGAGTCACGCTACAACTAATTACATTTCT	<u>161</u>	32			
USH2A-9F	TTCCTAGAGCCATACAGATACTTG	<u>162</u>	24	1826	9F/9R	21
USH2A-9R	GCTGAATGGAAACGGATGCTATT	<u>163</u>	23			